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Sequence 3, Application US/08977767

Sequence 3, Application US/08977767

Patent INFORMATION:
APPLICANT: Bandman, Olga
APPLICANT: Greenwald, Sara
APPLICANT: Corley, Nell C.
TITLE OF INVENTION: CARBONIC ANHYDRASE VIII
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTER READABLE FORM:
MEDIUM TYRE: DISKELE
COMPUTER ID SISKELE
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: DISKELE
COMPUTER: DISKELE
COMPUTER: DISKELE
COMPUTER: DISKER

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LENGTH: 1345 amino acids
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STRANDEDNESS: single
TOPOLOGY: linear
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; LIBRARY: GenBank
; CLONE: 1532042
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Sequence 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-07-906-349A-6
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US-08-244-941-2
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US-08-083-590A-19
US-08-532-384-19
US-08-185-432-18
US-08-144-121-3
US-08-735-893-3
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                         OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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Match Length DB
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Perfect score:
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Sequence 19, Appl
Sequence 19, Appl
Patent No. 5472939
Patent No. 525642
Patent No. 5472939
Patent No. 5472939
Patent No. 5472939
Patent No. 5472939
Sequence 23, Appl
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Sequence 24, Appl
Sequence 27, Appl
Sequence 6, Appl
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ALIGNMENTS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 476 AAAACTTACCTATAAATTAAGAAGAGTTTTTATTACAAATTTCAACCT------AAA 526
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SSRCKACPANKVQGAVATAGGTATLIA----QCALECPAGTVLTDGTTSTYKQAASECV 404
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                                                                                                                                                                                                                                                                                   253 AGCCTGTGAGGGAGTGCAGAACACCGTTTTC-----TTAATTTGGGTGTTCGACG 302
                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                          21 ANCPVGT-----ETNTAGQV-DDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKK 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   190 RLNFYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDA---TITAQCNVACPDGTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      247 SAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAGGAAT---
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                                                                                                                                                                                                    Indels 133;
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Sequence 11, Application US/08209521
Patent No. 5922855
GENERAL INFORMATION:
APPLICANT: Liskay, Robert M.
APPLICANT: Bronner, C. Eric
APPLICANT: Baker, Sean M.
APPLICANT: Rollag, Roni J.
APPLICANT: Rollag, Roni J.
TITLE OF INVENTION: MAMMALIAN DNA MISMATCH REPAIR GENES
WOMBER OF SEQUENCES: 30
                                                                                                                                                            Length 801;
                                                                                                                                                                                                                                                                                                                                                                        303 GAGACTGTCGTCGGACACTCCCTCACGTCTTGTGGTGGCCGGGTCACCCT--
                                                                                                                                                                                                                                                                                                                                                                                                                                                             353 GTGACCCTCTCACTTTGG--TTGGA-----ACTT-TAGGGG-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/209,521
                                                                                                                                                 Score 187.5; DB 1;
Pred. No. 2.3e-08;
                                                                                                                                                                                                           7; Mismatches
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                                                                                                                                                                 Query Match 7.4%;
Best Local Similarity 25.7%;
Matches 116; Conservative
801 amino acids
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MEDIUM TYPE: Floppy
                                                                                  protein
               TYPE: amino acid
STRANDEDNESS: sing
TOPOLOGY: linear
MOLECULE TYPE: prote
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 3
US-08-209-521-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                         US-07-906-349A-6
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    LENGIH:
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APPLICANT: Schlessinger, Joseph
APPLICANT: Margolis, Benjamin L.
TITLE OF INVENTION: IDENTIFYING TARGET PROTEINS FOR EUKARYOTIC TYROSINE KINASES ITTLE OF INVENTION: TARGET PROTEINS
TITLE OF INVENTION: TARGET PROTEINS
TITLE OF INVENTION: TARGET PROTEINS
TITLE OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                   17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         927 CITIGGCITCAICA-CCIAICAGGCICICICAGCACCCCAC--CCIGCI--CGGAGAC 980
                                                                                                                                                                                                                                                                                                                                                 190 RLNFYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVACPDGTISAA 249
                                                                                                                                                                                                                                                                                                                                                                                                                                      250 GVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAGGAATLAKQCN 309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   851 C-------AATGA-GATGC 882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 883 TAC------GITTCTTCAAGACCTGAGCCTGGAGCTCCTGTGCCCCGAGTC 926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          366 ATAGGTATLIAQCALECPAGTVLFDGTTSTYKQAASECVKCAANFYTTKQTDWVAGIDTC 425
                                                                                                                                                                                                                                                                 132 AGASICTACPVNRVGGALIAGNAAT--IVAQCNVACPIGTALDDGVTTDYVRSFTECVKC 189
                                                                                                                                 649 AGCACGGACGGAGCTGGGCTCTGAACACCAGGATCAACCATCAGGGTTTCTCTGTGAGGTG 708
                                                                                                                                                                              77 AQPNPPATANLVTQCNVKCPAG-----TAIAGGATDYAAIITECVNCRINFYNENAPNFN 131
                                                     Indels 103; Gaps
                                                                                           21 ANCPVGTETNTAGQVDDLG-TPANCVNCQKNFYYNNAAAFVPGAS---TCTPCPQKKDAG 76
                                                                                                                                                                                                                                                                                                         760 AGGGGCCCCAA--TGGCCTGGCCATTCTCAGCCTCTTTGTCAATGT------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         310 IACPDGTAIASGATNYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKV----QGAV
                                                                                                                                                                                                                                                                                                                                                                                          -----GGCTG-----GTAGCT-CAAACCCGTTCCTCAGACGCCTCCTTAACCGTGACA
           Length 1345;
                                                                                                                                                                                                           CURRENT APPLICATION OF FILLS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: PAPLICATION NUMBER: US/07/906,349A FILING DATE: 30-JUN-1992
               DB 2;
                                                       5; Mismatches 212;
           Score 200.5; DB 2
Pred. No. 3.3e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Browdy and Neimark
STREET: 419 Seventh Street, N.W.
CIIY: Washington
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07/643,237
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/643,23
FILING DATE: 18-JAN-1991
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                    7.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                    Query Match 7.9%
Best Local Similarity 25.8%
Matches 111; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            426 TSCNKKLTSGA 436
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US-07-906-349A-6
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                                                                                                                                                                                                                                                                                                                                                                                                         804
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TYPE: amino acid
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US-08-460-309-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       194 YYNGNNGNTPFNPGKSQCT-PCPAIKPANVAQATLGNDATI------TAQCNV 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                240 ACPDGTISAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAG 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              131 GCCAGTATTTCTACCTATGGCTTTCGAGGTGAGGTAAGGTT------TCAA 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           181 GAAATGTKTAAAATATCCTCCTGTGATGACATTGTXTGTCATTTGTTAGTATTTCT 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             48 -----TITITGTATCTAACAAAGAAGATCTGGATATTGTATGTAAGGKTCAC-T 99
                                                                                                                                                                                                                                                                                                                                                                                                  Ouery Match
7.1%; Score 179.5; DB 2; Length 341;
Best Local Similarity 25.8%; Pred. No. 3.8e-08;
Matches 98; Conservative 9; Mismatches 130; Indels 143;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 5, Application US/08460309
Patent No. 5837496
Patent No. 5837496
APPLICANT: Engvall, Eva
APPLICANT: Letvo, Ilmo
TITLE OF INVENTION: Nucleic Acids Encoding Merosin
TITLE OF INVENTION: Pragments and Uses Thereof
NUMBER OF SEQUENCE: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ....resSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
STATE: California
COUNTRY: USA
            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: VAR RYSSEABERGHE, PIETE C.
RECISTRATION NUMBER: 33,557
REFERENCE/DOCKET NUMBER: 045U 306A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (503) 224-6655
TELEFRAX: (503) 224-6655
TELEFRAX: (503) 235-6679
INFORMATION FOR SEO ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 341 amino acids
TYPE: amino acid
STRANBENESS: single
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FILING DATE: 08-MAR-1994
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MOLECULE TYPE: CDNA
POSITION IN GENOME:
MAP POSITION: 3P21.3-23
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                            US-08-209-521-11
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987 FY----AYODGS--CIPCDCPHIQNICDPEIG-----ECV--CPPHIQGVKCEECEDGH 1032
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        934 QCLHGYYGLDSGHG-----CRPCN--CSVAGSVSDGCTDEGQCHCVPGVAGKRCDRCAHG 986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    749 ------AECNVH---GVCI---ACAHNTTGVHČEQČLPGFYGEPSRGTPGDCQPC 791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            881 -----HCERCADGFYGDAVTAKNCRACECHVKGSHSAVCHLETGLCDCKPNVTGQQCD 933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           292 -----YGAEATAGGAATLAKQCNIACPDGTAIASGATNY-----VILQTECLNCAAN 338
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ouery Match
6.9%; Score 175.5; DB 2; Length 3075;
Best Local Similarity 21.2%; Pred. No. 1.6e-06;
Matches 110; Conservative 44; Mismatches 187; Indels 177; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18 IKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAFVPGASTCTPCPQKKDAGA 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             |||:
|792 -ACPL-----TIAS--NNFSPT-CHINDGDEVVCDWCAPGYSGAWCERCADG
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Ralease #1.0, Version #1.25
GURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NKKLTSGAEANLPES------AKKNI---QCD 451
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                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/125,077
FILING DATE: 22-SEP-1993
APPLICATION NUMBER: US PCT/US 94/10730
FILING DATE: 21-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/472,319
FILING DATE: 30-JAN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/919,951
FILING DATE: 27-JUL-1992
ATTOMEY/AGENT INFORMATION:
ANNUEL CAMPAINT OF THE PRIOR OF THE PRIOR DATE:
ANNUEL CAMPAINT OF THE PRIOR OF THE PRIOR DATE:
ANNUEL CAMPAINT OF THE PRIOR OF THE PRIOR DATE:
ANNUEL CAMPAINT OF THE PRIOR DATE:
AND DAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Campbell, Cathryn A.
REGISTRATTON NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LA 9721
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 3075 amino acids
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ADDRESSEE: COOLEY GODWARD CASTRO HUDDLESON & TATUM STREET: FIVE PALO ALTO SQUARE CITY: PALO ALTO CITY: CALIFORNIA COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/284,941
FILING DATE: 2 August 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6.6%; Score 166.5; DB 2; 20.7%; Pred. No. 2.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        53; Mismatches 135;
                                                                                                                                                                                                                                                                                                                                                                      1093 DLRGTSGDACNLEQGLCGCVEETGACPCKENVFGPQCN 1130
                                                                                                                                                                                                                                                                                                                            429 NKKLTSGAEANLPES-----AKKNI---QCD 451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: NEELEY PH.D., RICHARD L.
REGIESTRATION NUMBER: 30092
REFERENCE/DOCKET NUMBER: CHIR-009/01US
TELECOMMUNICATION INFORMATION:
TELEPAX: (415) 843-5070
TELEPAX: (415) 843-5070
TELEPAX: (415) 843-5070
SECURATION FOR SEQ. ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                        393 TSTYKQAASECVKCAANFYTTKQTDWVAGIDTCTS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/08284941
Patent No. 5863756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 6.6%
Best Local Similarity 20.7%
Matches 86; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 2 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94306
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                       881
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          78 OPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTC 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          138 TACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDG--VTTDYVR---SFTECVKCRLN 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    792 -ACPL------TIAS--NNFSPT-CHLNDGDEVVCDWCAPGYSGAWCERCADG 834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       253 NWVAQNTECTNCAPNFYNN--NAPN-------FNPGNSTCLPCPANKD-- 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      706 VEHCECPOGY-----TGTSCESCLSGYRRVDGILF---GGICQPCECHGHA-- 748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             749 -----ACAHNITGVHCEQCLPGFYGEPSRGTPGDCQPC 791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                            Sequence 5, Application US/08125077

Sequence 5872231

Patent No. 5872231

APPLICANT: Engvall. Eva APPLICANT: Leivo, Ilmo

TITLE OF INVENTION: Nucleic Acids Encoding Merosin, Merosin, TITLE OF INVENTION: Pragments and Uses Thereof

NUMBER OF SEQUENCES: 23

CORRESPONDENCE ADDRESS:

ADDRESSEE: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels 177;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6.9%; Score 175.5; DB 2;
21.2%; Pred. No. 1.6e-06;
tive 44; Mismatches 187;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/125,077
FILING DATE: 22-SEP-1993
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION UNDER: US PCT/US 94/10730
APPLICATION UNDER: 21-5EP-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LA 9721
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 07/472,319
FILING DATE: 30-JAN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/919,951
FILING DATE: 27-UUL-1992
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIDLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 3075 amino acids
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Best Local Similarity 21.24
Matches 110; Conservative
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California
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STATE:
                                                            US-08-125-077-5
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                                                                                                                                                                                                                                                                                           1033 WGYDAEVGCQACNCSLVGSTHHRCDVVTGHCQCKSKFGGRACVQCSLGYRDFPDCVPCDC 1092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        70 PQKKDAGAQPNPPATANLV--TQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENA 127
                                                                                                                                                                        987 FY----AYQDGS--CIPCDCPHIQNICDPEIG-----ECV--CPPHIQGVKCEECEDGH 1032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels 141; Gaps
                                                                                                                   339 FYFDGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQCALECPAGT-----VLTDGT 392
                                           934 QCLHGYYGLDSGHG-----CRPCN--CSVAGSVSDGCTDEGQCHCVPGVAGKRCDRCAHG 986
292 -----YGAEATAGGAATLAKQCNIACPDGTAIASGATNY-----VILQTECLNCAAN 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 969;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: BARR, PHILLP J
APPLICANT: KIEPER, MICHAEL C
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PACE 4 AND
TITLE OF INVENTION: PACE 4.1 GENE AND POLYPEPTIDES IN CELLS
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
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GENERAL INFORMATION:
APPLICANT: Barr, Philip J
APPLICANT: Riefer, Michael C
APPLICANT: Riefer, Michael C
TITLE OF INVENTION: Compositions and Methods for PACE 4 and 4.1 Gene and
TITLE OF INVENTION: Polypeptides in Cells
FILE REFERENCE: CHIR*009/04US
CURRENT APPLICATION NUMBER: US/09/236,503
CURRENT FILING DATE: 1999-01-25
EARLIER APPLICATION NUMBER: 08/447,642
EARLIER FILING DATE: 1994-08-02
EARLIER FILING DATE: 1994-08-02
EARLIER FILING DATE: 1994-08-02
EARLIER FILING DATE: 1994-08-02
EARLIER FILING DATE: 1992-03-09
NUMBER OF SEQ ID NOS: 16
                                                    721 GSVKTSRKCVSVCPLGYFG----DTAARRCRRCHKGCETCSS------RAAIQCL 765
                                                                                                                                                                                           KCRLNFYYNGNNGNTPFNPGKSQC-TPCPAIKPANVAQATLGNDATITAQCNVACPDGTI 246
                      PQKKDAGAQPNPPATANLV--TQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENA 127
                                                                                                      PNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECV 187
                                                                                                                                                                                                                                                                             SAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLP-CPANKDYGAEATAGGAATLA 305
                                                                                                                                                                                                                                                                                                                   -----KKCVDEPEKCTVC-----KEGFSLARGSCIPDCEPGTYFDSELIRCG----- 846
                                                                                                                                                                                                                                                                                                                                                                                          RCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFY 411
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KCRLNFYYNGNNGNTPFNPGKSQC-TPCPAIKPANVAQATLGNDATITAQCNVACPDGTI 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 128 PNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECV 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            247 SAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLP-CPANKDYGABATAGGAATLA 305
                                                                                                                                                                                                                                                                                                                                                                 306 KOCNIACPDGTAIASGATNYVILQTECLNCAANFYFD------GNNFQAGS-----S
                                                                                                                                                                                                                        897 VCRRCDEN----SRNCSRCKTGF-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels 141;
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20.7%; Pred. No. 2.1e-06;
Live 53; Mismatches 135;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 2, Application US/09236503
Patent No. 6277590
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Best Local Similarity 20.7%
Matches 86; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Ver. 2.0
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US-09-236-503-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-236-503-2
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                                           188 KCRLNFYYNGNNGNTPFNPGKSQC-TPCPAIKPANVAQATLGNDATITAQCNVACPDGTI 246
                                                                                                                             247 SAAGVINWVAQIITECTINCAPINFYININAPINFINFGINSTCLP-CPAIKDYGAEATAGGAATLA 305
                                                                                                                                                        352 RCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFY 411
                                                                                                                                                                                                                                                                                                                                                      -----CLSC-AG------SSRNCSRCKTGF- 922
                                                                                                                                                                                                               306 KQCNIACPDGTAIASGATNYVILQTECLNCAANFYFD------GNNFQAGS-----S
                                                                                  766 SCRRGFYHH-QEMNT-----CVTLCPAGFYADESQKN-----CLKCHPSC----
                                                                                                                                                                                                                                                        847 -ECHHTC--GTCVGPG-----REECIHCAKNFHFHDWKCVPACGEGFYPEEMPGLPHK
                                                                                                                                                                                                                                                                                                                                                                                      412 TTKQTDWVAGIDTCTSCNKKLT-SGAEANLPESAKKNIQCD---FANFLSISLLL 462
                                                                                                                                                                                                                                                                                                                                                                                                                               923 ------TQLGTSCITNHTCSNADETFCEMVKSNRLCERKLFIQFCCRTCLL 967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels 141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: BARR, PHILIP J
APPLICANT: KIEPER, MICHAELC
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PACE 4 AND
TITLE OF INVENTION: PACE 4.1 GENE AND POLYPEPTIDES IN CELLS
NUMBER OF SEQUENCES:
ADDRESSE: COOLEY GODMARD CASTRO HUDDLESON & TATUM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 969;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: COOLEY GODWARD CASTRO HUDDLESON & TATUM
STREET: FIVE PALO ALTO SQUARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/447,642

FILING DATE: 23-MAX-1995

CLASSIPICATION NUMBER: US 08/284,941

FILING DATE: 2 August 1994

ATTORNEY/AGENT INFORMATION:

NAME: NELLEY PH.D., RICHARD L.

REGISTRATION NUMBER: 30092

REFERENCE/DOCKET NUMBER: CHIR-009/01US

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEFAX: (415) 843-570

TELEFAX: (415) 843-567

TELECOMMUNICATION PRO SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:

LEMOTH: 969 mnino acids

LEMOTH: 969 mnino acids
721 GSVKTSRKCVSVCPLGYFG-----DTAARRCRRCHKGCETCSS--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2;
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6.6%; Score 166.5; DB 2;
Best Local Similarity 20.7%; Pred. No. 2.1e-06;
Matches 86; Conservative 53; Mismatches 135;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/08447642
Patent No. 5989890
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amino acid
                                                                                                                                                                                                                                                                                                                                            897 VCRRCDEN------
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; MOLECULE TYPE: protein US-08-447-642-2
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STATE: CALIFORNIA
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US-08-447-642-2
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897 VCRRCDEN-----
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STATE: California
COUNTRY: USA
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TOPOLOGY:
US-08-460-309-4
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US-08-460-309-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             70 POKKDAGAQPNPPATANLV--TQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENA 127
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                                                                                                                                                  897 VCRRCDEN-----SRNCSRCKTGF- 922
                                                                       306 KQCNIACPDGTAIASGAINYVILQTECLNCAANFYFD------GNNFQAGS----S 351
                                                                                                                   352 RCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFY
                                                                                                                                                                                       412 TTKQTDWVAGIDTCTSCNKKLT-SGAEANLPESAKKNIQCD---FANFLSISLLL 462
                                                                                                                                                                                                             Indels 141;
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                                                                                                                                                                                                                                                                                                      Sequence 2, Application PC/TUS9302147A
GENERAL INFORMATION:
APPLICANT: BARR, PHILIP J
APPLICANT: KIEFER, MICHAEL C
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PACE 4 AND
TITLE OF INVENTION: PACE 4.1 GENE AND POLYPEPTIDES IN CELLS
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOLEY GOWARD CASTRO HUDDLESON & TATUM
ADDRESSEE: COOLEY GODWARD CASTRO HUDDLESON & TATUM STREET: FIVE PALO ALTO SQUARE CITY: PALO ALTO STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SCHWARE: Patentin Release #1.0, Version #1.25 SCHRENT APPLICATION DATA:
APPLICATION NAMBER: PCT/US93/02147A FILING DATE: 1999309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 20.7%; Pred. No. 2.1e-06;
Matches 86; Conservative 53; Mismatches 135;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/648,629
FILING DATE: 09-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: NEELEY PH.D., RICHARD L
REGISTRATION NUMBER: 30092
REFERENCE/DOCKET NUMBER: CHIR-009/00US
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (415) 494-7622
TELEFAX: (415) 857-0663
TELEX: 380816 COOLEY PA
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 969 amino acids
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   805 ----KKCVDEPEKCTVC-
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94306
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PCT-US93-02147A-2
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PCT-US93-02147A-2
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                                                                                                                                                                                                                                                                                                      RCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFY 411
                                                                                                                                                                                                                                                                                                                                     188 KCRLNFYYNGNNGNTPFNPGKSQC-TPCPAIKPANVAQATLGNDATITAQCNVACPDGTI 246
                                                                                                  247 SAAGVINWVAQNTECTICAPNFYNINAPNFINFGINSTCLP-CPANKDYGAEATAGGAATLA 305
                                                                                                                                                                                                      -GNNFQAGS----S 351
                                                                                                                                                                                                                                                                                                                                                                                                          412 TTKQTDWVAGIDTCTSCNKKLT-SGAEANLPESAKKNIQCD---FANFLSISLLL 462
                                                                                                                                                                                                                                                                                                                                                                                                                                                          --TQLGTSCITNHTCSNADETFCEMVKSNRLCERKLFIQFCCRTCLL 967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Engvall, Eva
APPLICANT: Leivo, Ilmo
TITLE OF INVENTION: Nucleic Acids Encoding Merosin, Merosin
TITLE OF INVENTION: Fragments and Uses Thereof
CORRESPONDENCES: 23
CORRESPONDENCE ADDRESS:
                                                  766 SCRRGFYHH-QEMNT-----CVTLCPAGFYADESQKN-----CLKCHPSC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                             -----CLSC-AG-----
                                                                                                                                                                                                         306 KQCNIACPDGTAIASGATNYVILQTECLNCAANFYFD----
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FILING DATE: 22-SEP-1993
APPLICATION NUMBER: US PCT/US 94/10730
FILING DATE: 21-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/472,319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 07/472,319
FILING DATE: 30-JAN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/919,951
FILING DATE: 27-JUL-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P-LA 9721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4, Application US/08460309
Patent No. 5837496
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Campbell, Cathryn A. REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 4:
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LENGTH: 3111 amino aci
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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983 GQCWCQPGVTGKKCDRCAHGYF----NFQEGG--CTACECSHLGNNCDPKTG-----RC 1030
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                78 QPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRIN------FYNE 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    758 -----QC-----FGHAESCDDVTGECLNCKDHTGGPYCDKCLPGFYGE 795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            126 NAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTE 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CVKCRLNFYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCN-----V 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    240 ACPDGTISAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAG 299
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              379 ALECPAGTVLTDGTTSTYKQAASECVKCAANFY-------TTKQTDWVAGIDT 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 IKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGA 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    796 PTKGTSEDCQPC-ACPLN----IPSNNFS---PTCHLDRSLG-LICDGCPVGY--TGPR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          47; Mismatches 164; Indels 222;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 6.2%; Score 157; DB 2; Best Local Similarity 18.5%; Pred. No. 6.8e-05; Matches 98; Conservative 47; Mismatches 164
      PRICING DATE: 30-JAN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/919,951
FILING DATE: 27-JUL-1992
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LA 9721
TELECOMMUNICATION INFORMATION:
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERIZITICS:
LENGTH: 3111 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BANNER & ALLEGRETTI, LTD
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TITLE OF INVENTION: Laminin Chains:
TITLE OF INVENTION: Therapeutic Use
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: BANNER & ALLEGRETI, LJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15, Application US/08317450B Patent No. 5660982
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GENERAL INFORMATION:
APPLICANT: Tryggyason, Karl
APPLICANT: Kallunki, Pekka
APPLICANT: Pyke, Charles
                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 12
US-08-317-450B-15
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                                                                                                                                                                                                                                       78 OPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRIN------FYNE 125
                                                                                                                                                                                                                                                                                                  ------QC-----FGHAESCDDVTGECLNCKDHTGGPYCDKCLPGFYGE 795
                                                                                                                                                                                                                                                                                                                                                             126 NAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTE 185
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6.2%; Score 157; DB 2; Length 3111;
18.5%; Pred. No. 6.8e-05;
Ative 47; Mismatches 164; Indels 222; Gaps
                                                                                                                                                                   18 IKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAFFVPGASTCTPCPQKKDAGA 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          888 ICKPGT------TGRYCELCADGYFGDAVDAKN-----CQPCRCN-----AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1077 GOCNCHPKFSGAKCTECSRGHWNYPRCNLCDCFLPGTDATTCDSETKKCSC 1127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Engvall, Eva
APPLICANT: Leivo, Ilmo
TITLE OF INVENTION: Nucleic Acids Encoding Merosin TITLE OF INVENTION: Fragments and Uses Thereof
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSE: Campbell and Flores
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    425 ------CTSCNKK-----LTSGAEANLPESAKKNIQC 450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ----AATLAKQCNIACPDGTAIASGATNYV---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US PCT/US 94/10730
FILING DATE: 21-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/472,319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/125,077 FILING DATE: 22-SEP-1993 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4, Application US/08125077; Patent No. 587221
Patent No. 587221
; General Information:
                                                                98; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
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STATE: California
                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92122
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US-08-125-077-4
   Ouery Match
                                     Best Local
                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  186
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; Sequence 13, Application US/08317450B
                   CORRESPONDENCE ADDRESS:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             585 CRSDGT-----
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US-08-317-450B-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  460 KPCPCHN-----GFSCSVIPETEEVVCN-NCPPGV------TGARCELCADG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 5.9%; Score 150.5; DB 1; Length 1111; Best Local Similarity 20.0%; Pred. No. 6.5e-05; Matches 82; Conservative 49; Mismatches 138; Indels 141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               627 LQRWEALISKAQGGDGVVPDTELE------GRWQQAEQALQDILRDA 667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       358 ANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCA 407
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/317,450B
FILING DATE: 04-0CT-1994
CLASSIFICATION: 435
ATTONNEY/AGENT INFORMATION:
NAME: CLASO MATCH REGISTRATION NUMBER: 37,293
REFERENCE/POCKET UNBER: 94,778
TELECOMMULICATION INFORMATION:
TELECOMMULICATION: 1870-115-1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15, Application US/08800593
Patent No. 6143505
GENERAL INFORMATION:
APPLICANT: Tryggvason, Karl
APPLICANT: Kallunki, Pekka
APPLICANT: Pyke, Charles
APPLICANT: Trille OF INVENTION: Laminin Chains: Diagnostic and
TITLE OF INVENTION: Therapeutic Use
                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
  : Ten South Wacker Drive
Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 312-715-1234
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 1111 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; MOLECULE TYPE: protein US-08-317-450B-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
                                              Illinois
                                                                  USA
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                                                               COUNTRY:
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138 TACPVNRVGGALTAGNAATIVAQ-----CNVACPTGTALDDGVTTDYVRSFTECVKCRLN 192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                78 OPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTC 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       253 NWVAQNTECTNCAPNFYNNN-APNFNPGNSTCLPCPANKDYGAEATAGGAATLAKQCNIA 311
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5.9%; Score 150.5; DB 4; Length 1111;
Best Local Similarity 20.0%; Pred. No. 6.5e-05;
Matches 82; Conservative 49; Mismatches 138; Indels 141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             358 ANKVOGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCA 407
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ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive
CITY: Chicago
STATE: 1111nois
COUNTRY: USA
                                                                                                                                           ZIF: 60606

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRIT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                             CONTENT AFFLICATION NUMBER: US/08/800,593
FILING DATE: 18-FEB-1997
CLASSIPICATION: 435
PRIOR APPLICATION NUMBER: US 08/317,450
FILING DATE: 04-OCT-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Chao, Mark
REFERENCE/DOCKET NUMBER: 37,293
REFERENCE/DOCKET NUMBER: 94,778-B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LQRMEALISKAQGGDGVVPDTELE-
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TELEPHONE: 312-913-0001
TELEFAX: 312-913-0002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 15: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 1111 amino acids TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Score 150.5; DB 1; Length 1193;
; Pred. No. 7.1e-05;
49; Mismatches 138; Indels 141;
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                                                                                                                                                                                                                           COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/317,450B
FILING DATE: 04-OCT-1994
ATTORNEY AGENT
            APPLICANT: Trygovason, Karl
APPLICANT: Trygovason, Karl
APPLICANT: Kallunki, Pekka
APPLICANT: Pyke, Charles
TITLE OF INVENTION: Laminin Chains: Diagnostic and
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
                                                                                                                                                      E: BANNER & ALLEGRETTI, LTD.
Ten South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY AGENT INFORMATION:
NAME: Chao, Mark
REGISTRATION NUMBER: 37,293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 94
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-715-1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 312-715-1234
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARATERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.9%;
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Best Local Similarity 20.0%
Matches 82; Conservative
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                                                                                                                                                  ADDRESSEE: BANNES
STREET: Ten South
CITY: Chicago
STATE: Illinois
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        138 TACPVNRVGGALTAGNAATIVAQ-----CNVACPTGTALDDGVTTDYVRSFTECVKCRLN 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   460 KPCPCHN-----GFSCSVIPETEEVVCN-NCPPGV------TGARCELCADG 499
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  500 YFGDPFGEHGPVRP----CQPCQC--NSNVDPSASGNCDRLTGRC-LKCIHNT---AGI- 548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      253 NWVAQNTECTNCAPNFYNNN-APNFNPGNSTCLPCPANKDYGAEATAGGAATLAKQCNIA 311
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 1193;
                                                                                                           APPLICANT: Tryggvason, Karl
APPLICANT: Kallunki, Pekka
APPLICANT: Pyke, Charles
TITLE OF INVENTION: Laminin Chains: Diagnostic and
TITLE OF INVENTION: Therapeutic Use
                                                                                                                                                                                                                                                                     ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff STREET: 300 South Wacker Drive CITY: Chicago STATE: 1111nois COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
5.9%; Score 150.5; DB 4;
Best Local Similarity 20.0%; Pred. No. 7.1e-05;
Matches 82; Conservative 49; Mismatches 138;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/800,593
FILING DATE: 18-FEB-1997
CLASSIFICATION 435
PRIOR APPLICATION DATA:
APPLICATION UNBER: US 08/317,450
FILING DATE: 04-OCT-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94,778-B
                                             Sequence 13, Application US/08800593
Patent No. 6143505
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Chao, Mark
REGISTRATION NUMBER: 37,293
REFRENCE/POCKET NUMBER: 94,77
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 312-913-0002
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1193 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; MOLECULE TYPE: protein US-08-800-593-13
                                                                                                                                                                                                                             NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                          90909
RESULT 15
US-08-800-593-13
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585 CRSDGT
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Qy Dp

Search completed: December 26, 2001, 10:35:54 Job time: 145 sec.